

**REMARKS**

It is noted that this application is in condition for allowance except for formal matters. According to the first page of the action the allowed claims are claims 8-13. However, claims 6-13 are on file and were allowed previously. It is respectfully requested that the examiner confirm that claims 6-13 are allowable.

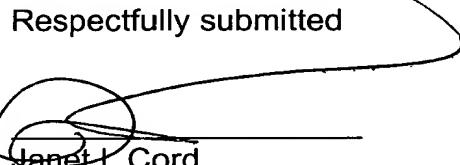
The specification has been amended to identify the sequences on pages 4, 16, 18 and 19 of the specification. A sequence listing and computer readable form is also being filed herewith.

A request for reconsideration of the decision dismissing the petition for acceptance of color photographs is being submitted herewith which includes an explanation of why color photographs are needed.

A new abstract of the disclosure is being submitted herewith.

It is respectfully requested that the notice of allowance be issued.

Respectfully submitted



Janet I. Cord  
c/o Ladas & Parry LLP  
26 West 61st Street  
New York, New York 10023  
Reg. No. 33, 778 (212-708-1935)



SEQUENCE LISTING

<110> KHANUJA, SUMAN PREET SINGH  
SHASANY, AJIT KUMAR  
DHAWAN, SUNITA  
DAROKAR, MAHENDRA PANDURANG  
SATAPATHY, SARITA  
KUMAR, TIRUPPADIRIPULIYUR R. SANTHA  
SAIKIA, DHARMENDRA  
PATRA, NIRMAL KUMAR  
BAHL, JANAK RAJ  
TRIPATHY, ARUN KUMAR  
KUMAR, SUSHIL

<120> NOVEL SCREENING METHOD FOR SELECTION OF INSECT TOLERANT PLANTS

<130> U-012567-2

<140> 09/487,405

<141> 2000-01-18

<160> 20

<170> PatentIn version 3.2

<210> 1

<211> 10

<212> DNA

<213> Artificial Sequence

<220>

<223> MAP Primer

<400> 1

aatcggagc

10

<210> 2

<211> 10

<212> DNA

<213> Artificial Sequence

<220>

<223> MAP Primer

<400> 2

gtcctactcg

10

<210> 3

<211> 10

<212> DNA

<213> Artificial Sequence

<220>

<223> MAP Primer

<400> 3

gtccttagcg

10

<210> 4

<211> 10

<212> DNA

<213> Artificial Sequence  
<220>  
<223> MAP Primer

<400> 4  
tgcgcgatcg 10

<210> 5  
<211> 10  
<212> DNA  
<213> Artificial Sequence  
<220>  
<223> MAP Primer

<400> 5  
aacgtacgca 10

<210> 6  
<211> 10  
<212> DNA  
<213> Artificial Sequence  
<220>  
<223> MAP Primer

<400> 6  
gcacgcccga 10

<210> 7  
<211> 10  
<212> DNA  
<213> Artificial Sequence  
<220>  
<223> MAP Primer

<400> 7  
caccctgcgc 10

<210> 8  
<211> 10  
<212> DNA  
<213> Artificial Sequence  
<220>  
<223> MAP Primer

<400> 8  
ctatcgccgc 10

<210> 9  
<211> 10  
<212> DNA  
<213> Artificial Sequence  
<220>  
<223> MAP Primer

<400> 9  
cgggatccgc 10

<210> 10  
<211> 10  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> MAP Primer

<400> 10  
gcgaattccg 10

<210> 11  
<211> 10  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> MAP Primer

<400> 11  
ccctgcaggc 10

<210> 12  
<211> 10  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> MAP Primer

<400> 12  
ccaagcttgc 10

<210> 13  
<211> 10  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> MAP Primer

<400> 13  
gtgcaatgag 10

<210> 14  
<211> 10  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> MAP Primer

<400> 14  
aggatacgtg 10

<210> 15  
<211> 10  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> MAP Primer

<400> 15  
aagatagcgg 10

<210> 16  
<211> 10  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> MAP Primer

<400> 16  
ggatctgaac 10

<210> 17  
<211> 10  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> MAP Primer

<400> 17  
ttgtctcagg 10

<210> 18  
<211> 10  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> MAP Primer

<400> 18  
catcccgAAC 10

<210> 19  
<211> 10  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> MAP Primer

<400> 19  
ggactccACG 10

<210> 20  
<211> 10  
<212> DNA

<213> Artificial Sequence

<220>

<223> MAP Primer

<400> 20

agcctgacgc 10